

IFWO

RAW SEQUENCE LISTING

DATE: 09/29/2004

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PATENT APPLICATION: US/10/815,337

TIME: 13:05:45

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73 Leu Ser Pro Ala Phe Gln Tyr Gly Asn Arg Thr Phe Thr Lys Tyr Pro

77 Glu Asp Ile Ser Asp Phe Phe Ile Gln Ser Phe Pro Ala Gly Phe Val

70

75

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85 Ser Asp Ile Asn Leu Ile Glu Glu Met Phe Val Tyr Arg Val Glu Tyr
86 115 120 125
89 Lys Gly Arg Asn Phe Pro Asn Asp Gly Pro Val Met Lys Lys Thr Ile
90 130 135 140
93 Thr Gly Leu Gln Pro Ser Phe Glu Val Val Tyr Met Asn Asp Gly Val
94 145 150 155 160
97 Leu Val Gly Gln Val Ile Leu Val Tyr Arg Leu Asn Ser Gly Lys Phe
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167 Gln Ile Arg Val Thr Lys Gly Ala Pro Leu Pro Phe Ala Phe Asp Ile
168 50 55 60 171 Leu Ser Pro Ala Phe Gln Tyr Gly Asn Arg Thr Phe Thr Lys Tyr Pro

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176	GIU	Asp	116	ser	85	PHE	Pite	116	G111	90	Pne	PIO	Ala	Grà	95	vai	
	Tyr	C1	7 ra	Thr		7,200	Tree	Clu	Λαn		C1**	T OIL	₹7-1	Clu		λνα	
180	тАт	GLU	Arg	100	neu-	Arg	1 Y L	GIU	105	GTÀ	Gry	nea	vaı	110	116	Arg	
	Cor	7 an	т 1 о		Lon	Tla	Clu	Clu		Dho	V-1	Ттт	7~~		C1.,	Tur	
	Ser	Asp	115	ASII	Leα	116	GIU	120	mec	File	val	TAT	125	var	GIU	ıyı	
184	T	C1		7. an	Dho	Dro	7 02		C1 11	Dro	7707	Mot		.T. r.ca	The	т1-	
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188	mb w		T 011	Cl.,	Drea	Cox	-	C111	₹7 - 1	77-7	TT rec	_	7 ~~	7 am	C1	17-3	
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	145 Leu	77-7	C1.,	Cln	Tro I	_	T 011	7727	m	7. ~~	_	7 an	Cor	Clar	Tara		
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223 225 227 229 231 233 235 237 249 241 248 249 251 253 254 257	acco gcct gagg ctgc atgt aaga ctgg atgc cagc <210 <211 <212 <213 <400 Met	ttggacategctagac	agg gaca to ca gaca to	gegteggeagegaegegaegegaegegaegegaegegae	ggtga gagco gagco gtggco gatco gaaga gacca ; 6 Ren: 6 Gln 5	aa ca ct gg cc cg ct ca gg cc ga gt ct gg ag ca ag ca illa	aacca gtgcat gcctt atcca ctggt cagca gtgta aaggg cacgt ctggg	acgton agaton consideration of the consideration of	g tto c cgo g tao c tto g ato c cto c cto g gto g cco nis	cacca cgtga cgcacca ccgaact cgagg gaaca gaagg cctgg	atgg acca acc geeg ageg tee gteg gact gget ggea	aggg aggg gcac gctt acat ccaa tgta gcaa tccc tcgt gcct	getge gegee ceate ceaac acta acta acta agtte agte ageac	Glu Val	caagg cctgo caagt cgago ccaco cgaco cagot ccact gcaco gtggg	ggcaac ccette caccc ggcacc gagggg gtgatg ggcgtg cgcac ctcatc gagacc gtgtaa	120 180 240 300 360 420 480 540 600
223 225 227 229 231 233 235 237 241 243 246 247 248 249 251 253 254 257 258	acco gcct gagg ctgc atgt aaga ctgg atgc cagca <210 <211 <212 <213 <400 Met 1 Ser	ttgga ttgtt tcga gcta gcta tcgt agac ttgga tcgc tcgc	agg gaca to ca	gegteggead gegadeggeeggeeggeggeggeggeggeggeggeggeggegge	ggtga gagco gattot ggtga ggtga gaaga gaaga gaaga gaaga gacca S	aa ca ct gg cc cg ct ca gg cc ga gt ct gg ag ca ac ct ag co	gecttateca gecttateca etggt cacaa cage gtgta aagge cacgt tetgge	acgte agate ccase agage ccage accge gcgte gcaae form Lys	g tto g tag g tag g tto g atc c cg g tto c ctg g gto g cco	cacca cgtga cggca cacct cgagg gaaca gaagg cctgg Thr 10 Val	atgg acca acc gccg agcg tcc gtcg gact ggct gg	aggg aggg gcac gctt acat ccaa tgta gcaa tccc tcgt gcct	getge gegee ceae ceae acate acate agtte cegae cagea	Glu Val	caage catgo caage gatco cace cace gatgo Ile 15 Phe	ggcaac ccette caccc ggcacc gagggg gtgatg ggcgtg cgcac ctcatc gagacc gtgtaa Met	120 180 240 300 360 420 480 540 600
223 225 227 229 231 233 235 237 241 243 246 247 248 249 251 253 254 257 258	acco gcct gagg ctgc atgt aaga ctgg atgc cagc <210 <211 <212 <213 <400 Met	ttgga ttgtt tcga gcta gcta tcgt agac ttgga tcgc tcgc	agg gaca to ca	gegteggead gegadeggeeggeeggeggeggeggeggeggeggeggeggegge	ggtga gagco gattot ggggg ggtga gaaga gaaga gaaga gacca S9 Ren: 6 Gln 5	aa ca ct gg cc cg ct ca gg cc ga gt ct gg ag ca ac ct ag co	gecttateca gecttateca etggt cacaa cage gtgta aagge cacgt tetgge	acgte agate cccae agage ccage accge accge gcgte gcaae form	g tto g tag g tag g tto g atc c cg g tto c ctg g gto g cco	cacca cgtga cggca cacct cgagg gaaca gaagg cctgg Thr 10 Val	atgg acca acc gccg agcg tcc gtcg gact ggct gg	aggg aggg gcac gctt acat ccaa tgta gcaa tccc tcgt gcct	getge gegee ceae ceae acate acate agtte cegae cagea	Glu Val	caage catgo caage gatco cace cace gatgo Ile 15 Phe	ggcaac ccette caccc ggcacc gagggg gtgatg ggcgtg cgcac ctcatc gagacc gtgtaa Met	120 180 240 300 360 420 480 540 600

RAW SEQUENCE LISTING DATE: 09/29/2004
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265 Gln Ile Arg Val Thr Lys Gly Ala Pro Leu Pro Phe Ala Phe Asp Ile
269 Leu Ser Pro Ala Phe Gln Tyr Gly Asn Arg Thr Phe Thr Lys Tyr Pro
270 65
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273 Glu Asp Ile Ser Asp Phe Phe Ile Gln Ser Phe Pro Ala Gly Phe Val
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277 Tyr Glu Arg Thr Leu Arg Tyr Glu Asp Gly Gly Leu Val Glu Ile Arg
278
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281 Ser Asp Ile Asn Leu Ile Glu Gly Met Phe Val Tyr Arg Val Glu Tyr
282
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285 Lys Gly Arg Asn Phe Pro Asn Asp Gly Pro Val Met Lys Lys Thr Ile
286
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                                                140
289 Thr Gly Leu Gln Pro Ser Phe Glu Val Val Tyr Met Asn Asp Gly Val
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                                             155
293 Leu Val Gly Gln Val Ile Leu Val Tyr Arg Leu Asn Ser Gly Lys Phe
                    165
                                        170
297 Tyr Ser Cys His Met Arg Thr Leu Met Lys Ser Lys Gly Val Val Lys
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301 Asp Phe Pro Glu Tyr His Phe Ile Gln His Arg Leu Glu Lys Thr Tyr
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305 Val Glu Asp Gly Gly Phe Val Glu Gln His Glu Thr Ala Ile Ala Gln
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309 Leu Thr Ser Leu Gly Lys Pro Leu Gly Ser Leu His Glu Trp Val
310 225
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314 <211> LENGTH: 720
315 <212> TYPE: DNA
316 <213> ORGANISM: Renilla reniformis
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321 aacctggagg gegtggtgaa caaccaegtg ttcaccatgg agggctgegg caagggcaac
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323 atcetgtteg geaaceaget ggtgeagate egegtgacea agggegeece eetgeeette
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325 geettegaea teetgageee egeetteeag taeggeaace geacetteae caagtaeeee
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327 gaggacatea gegaettett catecagage tteecegeeg gettegtgta egagegeace
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329 atgcgctacq aggacggcgg cctggtggag atccgcagcg acatcaacct gatcgaggag
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331 atgttegtgt acegegtgga gtacaaggge egeaacttee ceaacgaegg eccegtgatg
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333 aagaagacca tcaccggcct gcagcccagc ttcgaggtgg tgtacatgaa cgacggcgtg
                                                                          480
335 etggtgggee aggtgateet ggtgtaeege etgaacageg geaagtteta eagetgeeae
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337 atgcgcaccc tgatgaagag caagggcgtg gtgaaggact tccccgagta ccacttcatc
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339 cagcaccgcc tggagaagac ctacgtggag gacggcggct tcgtggagca gcacgagacc
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341 gecategece agetgaceag cetgggeaag cecetgggea geetgeaega gtgggtgtaa
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345 <211> LENGTH: 239
346 <212> TYPE: PRT
347 <213> ORGANISM: Renilla reniformis
349 <400> SEQUENCE: 8
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355 Ser Phe Lys Val Asn Leu Glu Gly Val Val Asn Asn His Val Phe Thr
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359 Met Glu Gly Cys Gly Lys Gly Asn Ile Leu Phe Gly Asn Gln Leu Val
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363 Gln Ile Arg Val Thr Lys Gly Ala Pro Leu Pro Phe Ala Phe Asp Ile
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367 Leu Ser Pro Ala Phe Gln Tyr Gly Asn Arg Thr Phe Thr Lys Tyr Pro
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371 Glu Asp Ile Ser Asp Phe Phe Ile Gln Ser Phe Pro Ala Gly Phe Val
372
375 Tyr Glu Arg Thr Met Arg Tyr Glu Asp Gly Gly Leu Val Glu Ile Arg
376
379 Ser Asp Ile Asn Leu Ile Glu Glu Met Phe Val Tyr Arg Val Glu Tyr
380
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383 Lys Gly Arg Asn Phe Pro Asn Asp Gly Pro Val Met Lys Lys Thr Ile
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387 Thr Gly Leu Gln Pro Ser Phe Glu Val Val Tyr Met Asn Asp Gly Val
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391 Leu Val Gly Gln Val Ile Leu Val Tyr Arg Leu Asn Ser Gly Lys Phe
392
                    165
                                         170
395 Tyr Ser Cys His Met Arg Thr Leu Met Lys Ser Lys Gly Val Val Lys
                                    185
                180
399 Asp Phe Pro Glu Tyr His Phe Ile Gln His Arg Leu Glu Lys Thr Tyr
            195
                                200
403 Val Glu Asp Gly Gly Phe Val Glu Gln His Glu Thr Ala Ile Ala Gln
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407 Leu Thr Ser Leu Gly Lys Pro Leu Gly Ser Leu His Glu Trp Val
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412 <211> LENGTH: 720
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414 <213> ORGANISM: Renilla reniformis
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419 aacctggagg gcgtggtgaa caaccacgtg ttcaccatgg agggctgcgg caagggcaac
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421 atcctgtccg gcaaccagct ggtgcagatc cgcgtgacca agggcgcccc cctgcccttc
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423 geettegaca teetgageee egeetteeag taeggeaace geacetteae caagtaeeee
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425 qaqqacatca qcqacttctt catccaqaqc ttccccqccq qcttcqtqta cqaqcqcacc
                                                                           300
427 etgegetaeg aggaeggegg eetggtggag ateegeageg acateaacet gategaggag
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429 atgttegtgt acegegtgga gtacaaggge egeaacttee ceaacgaegg ceeegtgatg
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431 aagaagacca tcaccggcct gcagcccagc ttcgaggtgg tgtacatgaa cgacggcgtg
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433 etggtgggee aggtgateet ggtgtaeege etgaacageg geaagtteta cagetgeeae
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435 atgcgcaccc tgatgaagag caagggcgtq gtgaaggact tccccgagta ccacttcatc
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437 cagcaccgcc tggagaagac ctacgtggag gacggcggct tcgtggagca gcacgagacc
439 gccatcgccc agctgaccag cctgggcaag cccctgggca gcctgcacga gtgggtgtaa
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443 <211> LENGTH: 239
444 <212> TYPE: PRT
445 <213> ORGANISM: Renilla reniformis
447 <400> SEQUENCE: 10
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RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 09/29/2004

PATENT APPLICATION: US/10/815,337

TIME: 13:05:46

Input Set : A:\Sequence listing.ST25.txt Output Set: N:\CRF4\09292004\J815337.raw

nvalid <213> Response:

se of "Artificial" only as "<213> Organism" response is incomplete, er 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

eq#:63,64,65,66,67,68,70,71,72,73,74,75,76,77,78,80

VERIFICATION SUMMARY .

DATE: 09/29/2004

PATENT APPLICATION: US/10/815,337 TIME: 13:05:46

Input Set : A:\Sequence listing.ST25.txt Output Set: N:\CRF4\09292004\J815337.raw

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date